```
48kD i-antigen pro
48 kDa immobilizat
IAG48 (G1) surface
55kD i-antigen pro
55 kDa immobilizat
Synthetic 55kD i-a
48kD i-antigen re
48kD i-antigen re
48kD i-antigen re
                                                                                                                February 11, 2003, 19:43:19 ; Search time 32.0571 Seconds (without alignments) 1837.243 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1: /SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980_DAT:*
2: /SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981_DAT:*
3: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
4: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
5: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
5: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
6: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
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8: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
8: /SIDS2/gcgdata/geneseqy-embl/AA1981_DAT:*
8: /SIDS2/gcgdata/geneseqy-embl/AA1991_DAT:*
8: /SIDS2/gcgdata/geneseqy-embl/AA2000_DAT:*
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8: /SIDS2/gcgdata/geneseqy-embl/AA2001_DAT:*
                                                                                                                                                                                                   US-09-497-967-6
2342
1 MKYNILLILIISLFINELRA.....STTFAKFLSISLLFISFYLL
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                        of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                  908470 seqs, 133250620 residues
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    protein search, using sw model

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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB25859
AAY97176
AAB25869
AAY97177
AAY97177
AAB25862
AAB25862
AAB25863
AAB25861
                                                                                                                                                                                                                                                                                     BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A_Geneseq_101002:*
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length: 2000000000
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93.4
39.3
39.3
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20.2
119.3
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Perfect score:
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Maximum DB seq
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SSKD I-antigen ami Human presentiin e Mouse laminin-15 a Wouse laminin-15 a VspA6-Si gene prod G protein-coupled Mouse laminin A ch Mouse laminin-1 al Drosophila melanog Human laminin 2 ma Human laminin 2 ma Merosin major subu Human laminin 2 ma Merosin major subu Human laminin 2 al Protein LAMA2 diff Novel human diagno Novel human diagno 1-aminocyclopropan Human laminin alph Human laminin alph Human laminin A ch 55kD 1-antigen ami Human laminin alph Sequence of mouse Human laminin alph Mouse laminin 2 am Mouse laminin 3 an Mouse laminin Gl C Drosophila melanog AAB18144 AAB25888 AAO14246 ABB81589 AAM50357 AAB25890 AAU07370 AAE11215 ABB64261 AAO17359 AAB19794 AAB19792 AAR71730 AAU07343 AAE17310 AAB25885 ABB81598 AAP94758 ABB81588 AAB19791 AAB19793 AAE17309 AAW50892 ABB71543 AAY15458 AAB19796 AAW50897 15 69 175 88 175 88 175 89 185 89 185 89 185 89 186 376 218.5 2018.5 202.5 202.5 202.5 199.5 198.1 191.5 191.1 1 

ALIGNMENTS

Immobilisation antigen; i-antigen; ichthyophthiriaais; vaccine; white spot disease; freshwater fish; immune response; infection control. AAB25859 standard; Protein; 442 AA FOUND INC 48kD i-antigen protein sequence. ) UNIV GEORGIA RES FOUND CORNELL RES FOUND INC. CLARK T G. DICKERSON H W. Chthyophthirius multifiliis 99US-0118634. 99US-0122372. 99US-0124905. 99US-0131121. 04-FEB-2000; 2000WO-US02962. 18-DEC-2000 (first entry) WO200046373-A1. 04-FEB-1999; 02-MAR-1999; 17-MAR-1999; 27-APR-1999; 10-AUG-2000 (UYGE-) (DICK/) (CORR ) 

Lin T;

Dickerson HW,

Clark TG,

AAY97176;

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This invention relates to novel i-antigen polypeptide sequences.

C I-antigens or immobilisation antigens ar common to a variety of
   I-antigens or immobilisation antigens ar common to a variety of
   whenostomatid ciliates and their expression varies in response to
   environmental stimuli. This invention relates to i-antigens in
   controlled the causing inchtyophthiriasis or white spot disease. The
   configuration includes two polypeptide and polymorleotide sequences for two
   invention includes two polypeptide and polymorleotide sequences for two
   invention includes two polypeptide and polymorleotide sequences for two
   invention includes two polypeptide and polymorleotide sequences or including I multifilis serotypes using the nucleotide sequences.

C or identifying I multifilis serotypes using the nucleotide sequences.

C an immune response in fish is useful for prophylaxis, treatment or for an immune response in fish is useful for prophylaxis, treatment or for controlling I multifilis infection in fish. Polymodicotide or protein
   controlling I multifilis infection in fish. Sequences AAA97042,
   controlling I multifilis infection in fish. Sequences AAA97042,
   controlling I multifilis infection in fish. Sequences AAA97043-A97044,
   cragments identified in the invention. Sequences AAA97043-A97064
   cradments identified in the sequences.

C scolation of the invention. Sequences AAB97043-A97064
   cradments identified in the sequences.

C scolation of the invention. Sequences AAB970843-B999 and
   cradments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVYAEATQKVQ 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      241 AAGAAAAGVAAVTSQCVPCQINKNDSPATAGAQANLATQCSTQCPTGTAIQDGVTLVFSN 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGAQANLATQCSNQCPTGTVLDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVF 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNKNDSPAT 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EANGNOPFAANNAARGICVPCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVTDVF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SSTQCSQCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTVLD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      isolation of the i-antigen gene sequences. Sequences AAB25859-B256
AAB25893-B25906 represent i-antigen protein and peptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Score 2342; DB 21;
; Pred. No. 2.2e-171;
0; Mismatches 0;
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AAY97176
ID AAY97176 standard; Protein; 442 AA.
                                                                                                                                       Claim 1; Figure 1; 144pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CASTTFAKFLSISLLFISFYLL 442
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Best Local Similarity 100.
Matches 442; Conservative
             WPI; 2000-506071/45.
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Gaps 9

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Length 442;

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Tetrahymena thermophila expresses two major beta-tubulin genes (BTU1 and BTU2), which encode identical beta-tubulin proteins. Either of these two BTU2), which encode identical beta-tubulin proteins. Either of these two genes (but not both at once) can be disrupted without a detectable change can be treas of the BTU1 beta-tubulin in the cell phenotype. A K150L substitution in the BTU1 beta-tubulin confers increased resistance to microtubule-stabilizing drug. Cc discaration of the BTU1-IK150M allele can be transformed to paclitaxel cc resistance by gene replacement of BTU1-IK150M with a wild-type BTU1 gene cc resistance by gene replacement of BTU1-IK150M with a wild-type BTU1 gene containing the bull-IK150M allele, BTU1: not a T. thermophila mutant constitutes the coding region of the neol gene (conferring resistance to substitutes the coding region of the neol gene (conferring resistance to parmomycin) for that of BTU1, can be used to generate BTU1 gene knockouts constitutes the coding region of Heterologous nucleic acids (especially concessful cell-surface expression that is maintained by way of negative successful cell-surface expression that is maintained by way of negative successful cell-surface expression that is maintained by way of negative concessful cell-surface expression that is maintained by way of negative successful recombination-mediated insertion of a heterologous nucleic constitution and immune response in a vertebrate. The transgenic constitution immune response in a vertebrate. The transgenic constitution in mune also useful for producing polytologulary. Tetrahymena expressing Ichthyophthius continued effective vehicles for vaccination of freshwater fish against are also vaccination of freshwater fish against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Recombinant expression systems for expressing heterologous nucleic acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                BTU1; beta-tubulin; protein expression system; negative selection; paclitaxel sensitivity; cell surface, antigen; protozoa; ciliate; live vaccine; Ichthyophthiius multifiliis; immobilization-antigen; i-antigen; freshwater; fish; protozoacide.
                                                                                                                                                                                                                                                                                            /note= "Gln encoded by CAR, TAG or TAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clark TG;
                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 3A; 83pp; English.
                                                                                        48 kDa immobilization-antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2000; 2000WO-US02966.
                                                                                                                                                                                                                           Ichthyophthiius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; AAA52134, AAA52135.
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                                                     (first entry)
                                                                                                                                                                                                                                                                                    Misc-difference 1..442
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DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000-514962/46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CLAR/) CLARK T G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAERTIG
                                                                                                                                                                                                                                                                                                                                           WO200046381-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-MAR-1999;
17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYGE-) UNIV
                                                     04-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                      10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GAER/)
(DICK/)
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442 AA;

Sequence

360

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55kD i-antigen protein of parasite isolate G5.
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                                                                                                                                                                                         409 AA;
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Extxxxxx
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                                                                                                                                                                                                                                                                                                           Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine;
white spot disease; freshwater fish; immune response; infection control.
                                                 AAGAAAAGVAAVTSQCVPCQINKNDSPATAGAQANLATQCSTQCPTGTAIQDGVTLVFSN 300
                                                                                                                                      Gaps
                                                                                                                      240
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                         1 MKYNILLILIISLFINELRAVPCPDGTQTQAGLTDVGAADLGTCVNCRPNFYYNGGAAQG
                                DRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNKNDSPAT
                                                                                  AGAQANLATQCSNQCPTGTVLDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVF
                                                                                                             SSTQCSQCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTVLD
                                                                                                                                                                                         DGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVYAEATQKVQ
    Length 442;
                Indels
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100.0%; Score 2342; DB 2.
100.0%; Pred. No. 2.2e-17.
ive 0; Mismatches 0
                                                                                                                                                                                                                                                                                              IAG48 (G1) surface protein amino acid sequence.
                                                                                                                                                                                                                                                       AAB25889 standard; Protein; 409 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                     FOUND INC
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                                                                                                                                                                                                                                                                                                                               Ichthyophthirius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                          99US-0122372.
99US-0124905.
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                                                                                                                                                                                                                                                                                                                                                                                   99US-0118634.
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             Conservative
                                                                                                                                                                                                                                                                                  (first entry)
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CLARK T G.
DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                  UNIV GEORGIA RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dickerson HW,
      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                            WO200046373-A1,
      Best Local Sim:
Matches 442;
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17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                                                  18-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                   04-FEB-1999;
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Query Match
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(CLAR/)
(DICK/)
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in controlling and their expression which is an oblique parasite of freshwater fish causing ichthyophthirias or white spot disease. The coff reshwater fish causing ichthyophthiriasis or white spot disease. The coff reshwater fish causing ichthyophthiriasis or white spot disease. The coff invention includes two polypeptide and polynucleotide sequences for two antigons, of 48 and 55 kD. Also included in the invention are controlling I multifillis serotypes using the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting on immune response in fish is useful for prophylaxis, treatment or for cortrolling I. multifillis infection in fish. Polynucleotide or protein antigenic i-antigen polypeptide obtained is also useful for treating or preventing I multifillis infection in fish. Sequences AAA97063, AAA97065 and AAAA97065 and AAAA97060, AAA97065 and AAAA97061 and AAA97061 and AAA9708 represent primers used in the container of an antigen genes and gene and containing in the invention. Sequences AAA9704 and in the container of an antigen genes and gene and containing in the invention. Sequences AAA97041 and gene and containing an antigen genes and gene and containing in the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (excluding AAA97060) and AAA97071-A9708B represent primers used in the isolation of the 1-antigen gene sequences. Sequences AAB25859-B25889 and AAB25893-B25906 represent 1-antigen protein and peptide sequences.
Novel 1-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  260 QINKNDSPATAGAQANLATQCSTQCPTGTAIQDGVTLVFSNSSTQCSQCIANYFFNGNFE
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                                                                                                                                                                     Disclosure; Figure 8; 144pp; English.
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I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
hymenostomatid ciliates and their expression varies in response to
controlmental stimuli This invention relates to i-antigens in
environmental stimuli This invention relates to i-antigens in
controlment fish causing ichthyophthiriasis or white spot disease. The
controlment includes two polypeptide and polynucleotide sequences for two
invention includes two polypeptide and polynucleotide sequences for two
controlling I multifilis sectul for the invention are
an immune response in fish is useful for prophylaxis; treatment or for
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controlling I. multifilis infection in fish. Sequences AAA37043-A97042,
controlling I. multifilis infection in fish. Sequences and gene
controlling I. multifilis infection in Sequences and Gene
controlling I. multifilis infection in Sequences AAA37035-A97042,
contigents identified in the invention. Sequences AAA37043-A97064
cectuding AAA37060) and AAA37089 represent i-antigen genes and gene
cectuding AAA37060) and AAA37089 represent primers used in the
cectuding AAA37060 and AAA37089 represent primers used in the
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cectuding AAA37060 represent i-antigen protein and peptide sequences.
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                      Immobilisation antigen; i-antigen; ichthyopthhiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    57 AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel i antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -------NAARGICVPCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         39.3%; Score 921; DB 21; Length 468; 41.8%; Pred. No. 1.9e-62; Live 45; Mismatches 139; Indels 11.
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                                                                                                                                                                                                                                                                                                                                                                      (UYGE-) UNIV GEORGIA RES FOUND INC. (CORR ) CORNELL RES FOUND INC. (CLAR/) CLARK T G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Figure 3; 144pp; English.
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                                                                                                                                                                                                                                                               99US-0118634.
99US-0122372.
99US-0124905.
                                                                                             Ichthyophthirius multifiliis.
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infection in fish
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                                                                                                                                                                                                                          04-FEB-2000;
                                                                                                                                                                                                                                                                                          02-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                    27-APR-1999;
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Best Local
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293 GVTLVFSNSSTQCSQCIANYFFNG-NFEAGKSQCLKCPVSKTTPAHA-PGNTATQATQCL 350
                                                                                                                                                                                                                                                               351 TICPAGTVLDDGTSINFVASATECTKCSAGFFASKTIGFTAGTDTCTECTKKLTSGATAK 410
                                                                                                                                                                                                                                                                                                                                          APGVQVEAAGAAAAGVAAVTSQCVPCQINKN-DSPATAGAQANLATQCSTQCPTGTAIQD 292
                                                                                                                                                                                              273 APN---FNPG-----NSTCLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIAS 320
DVFDRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--N 175
                                                                             176 DSPATAGAQANLATQCSNQCPTGTVLDDGVT--LVFNTSATLCVKCRPNFYYNGGSPQGE 233
                                                                                                        Recombinant expression systems for expressing heterologous nucleic acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BTU1; beta-tubulin; protein expression system; negative selection; paclitaxel sensitivity; cell surface; antigen; protozoa; ciliate; live vaccine; Ichthyophthius multifiliis; immobilization-antigen; i-antigen; freshwater; fish; protozoacide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Gln encoded by CAR or TAA"
                                                                                                                                                                                                                                                                                                                                                                                                       VYAEATQKVQCASTTFAKFLSISLLFISFYLL 442
                                                                                                                                                                                                                                                                                                                                                                                                                              Clark TG;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY97177 standard; Protein; 468 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    55 kDa immobilization antigen.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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DICKERSON
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17-MAR-1999;
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Immobilisation antigen; 1-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.

Ichthyophthirius multifillis.

Synthetic.

WO200046373-A1.

10-AUG-2000

Synthetic 55kD 1-antigen protein L6P.

18-DEC-2000 (first entry)

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Tetrahymena thermophila expresses two major beta-tubulin genes (BTU1 and BTU2), which encode identical beta-tubulin proteins. Either of these two genes (but not both at once) can be disrupted without a detectable change in the call phenotype. A #350L substitution in the BTU1 beta-tubulin can detectable change protein confers increased resistance to microtubule-depolymerizing drugs can dincreased sensitivity to paclitaxel, a microtubule-depolymerizing drugs calls carrying the Btu1-18750M allele. Earn beta means for positive containing the Btu1-18750M allele. Btu1-18750W with a wild-type BTU1 gene containing the Btu1-18750M allele. BTU1: neol construct, which containing the Btu1-18750M allele. BTU1: neol construct, which containing the Btu1-18750M allele. BTU1: neol construct, which parmonycin for that of BTU1, can be used to generate BTU1 gene knockouts concessful cell-surface expression be inserted into a BTU1 gene knockouts concessful cell-surface expression that is maintained by way of negative successful cell-surface expression that is maintained by way of negative concessful cell-surface expression that is maintained by way of negative concomponent response in a vertebrate. The transgenic containing resistance to paclitaxel in the resulting contained). In particular, Tetrahymena expression of the transgenic claimed). In particular, Tetrahymena expression of transgenic claimed). In particular, Tetrahymena expression of transgenic claimed). In particular, Tetrahymena expressing Ichthyophthius multifilis immobilization-antigen (1-antigen) protein on their surface are effection by I. multifillis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    117 NCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVT 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         118 DVFDRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--N 175
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             234 APGVQVFAAGAAAAGVAAVTSQCVPCQINKN-DSPATAGAQANLATQCSTQCPTGTAIQD 292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              39.3%; Score 921; DB 21; Length 468;
41.8%; Pred. No. 1.9e-62;
Live 45; Mismatches 139; Indels 114;
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(UYGE-) UNIV GEORGIA RES FOUND INC. (CORR ) CORNELL RES FOUND INC. (CLAR/) CLARK T G.

DICKERSON H W.

(CLAR/) (DICK/) (LINT/)

LIN T

99US-0122372.

99US-0131121

27-APR-1999;

99US-0118634

04-FEB-1999; 7-MAR-1999;

04-FEB-2000; 2000WO-US02962.

Lin

Dickerson HW,

Clark TG,

WPI; 2000-506071/45.

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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
Chymorostomatid cilitates and their expression varies in response to
Chymorothyphthitius multifillis, a protozoan which is an obligate parasite
of freshwater fish causing ichthyophthiriasis or white spot disease. The
invention includes two polypeptide and polymorlectide sequences for two
chitodies capable of binding to the nucleotide sequences and a method
controlling in multifilis serotypes using the nucleotide sequences.

A composition (containing the i-antigen nucleotide or papeling
controlling I. multifilis infection in fish. Polynucleotide or protein
vaccines comprising a portion of the amplified product encoding an
controlling I. multifiliis infection in fish. Sequences AAA97016-A97042.

Chymorothype in multifilis infection in fish. Sequences AAA97016-A97042.

Chymorothype in multifilis infection in fish. Sequences AAA97016-A97042.

Chragments identified in the invention. Sequences AAA97041-A97064

Chymorothype in multifilis infection in Sequences AAA97041-A97064

Chragments identified in the invention. Sequences AAA97041-A97064

Chymorothype in the invention of the invention of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel 1-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MKYNILLILIISLFINELRAVPCPDGTQTQ-AGLTDVGAADLGT---CVNCRPNFYYNGG 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the i-antigen gene sequences. Sequences AAB25859-B256
306 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels 114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 468;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        39.0%; Score 914; DB 21;
41.6%; Pred. No. 6.4e-62;
Live 45; Mismatches 140;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA------AGEANGNOPFAAN-------
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 213; Conserv
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AAB25882 standard; Protein; 468

RESULT 6 AAB25882 ID AAB

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WPI; 2000-506071/45.
                                                                                                                                                                         Sequence
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(CLAR/)
(DICK/)
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                                                                                                                                                                                                                                                                                         RESULT 8
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                                                                                                                                                                                                                                                                                                                       Immobilisation antigen; i-antigen; ichthyophthirlasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                      351 TTCPAGTVLDDGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAK 410
                                                                                                                                                                                   292
                                                                                                                          350
                                                                                    233
                                 117 NCRINFYNENAPNENAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVT 176
                                                     DVFDRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--N 175
                                                                DSPATAGAQANLATQCSNQCPTGTVLDDGVT - - LVFNTSATLCVKCRPNFYYNGGSPQGE
                                                                                                 222 VAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTE---CINCAPNFYNN-----N
                                                                                                                234 APGVQVFAAGAAAAGVAAVTSQCVPCQINKN-DSPATAGAQANLATQCSTQCPTGTALQD
                                                                                                                                             293 GVTLVFSNSSTQCSQCIANYFFNG-NFEAGKSQCLKCPVSKTTPAHA-PGNTATQATQCL
        AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV
                         -- NAARGICVPCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVT
                                                                                                                                                                                                                                                                                                          repeat amino acid sequence SEQ ID 9.
                                                                                                                                                                                                           VYAEATQKVQCASTTFAKFLSISLLFISFYLL 442
                                                                                                                                                                                                                    Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 UNIV GEORGIA RES FOUND INC.
CORNELL RES FOUND INC.
CLARK T G.
DICKERSON H W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lin T;
                                                                                                                                                                                                                                                                AAB25862 standard; Protein; 89
                                                                                                                                                                                                                                                                                                                                                                                                             99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                 Ichthyophthirius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                              04-FEB-2000; 2000WO-US02962.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HW,
                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dickerson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                              48kD i-antigen
                                                                                                                                                                                                                                                                                                                                                                 WO200046373-A1
                                                                                                                                                                                                                                                                                                                                                                                                              04-FEB-1999;
02-MAR-1999;
17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                                                                18-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clark TG,
                                                                                                                                                                                                                                                                                 AAB25862;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CORR )
(CLAR/)
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                                                                                                                                                                                                                                                    RESULT 7
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
the invitonmental stimul. This invention relates to i-antigens in
environmental stimul. This invention relates to i-antigens in
chithyophthirius multifilis, a protozoan which is an obligate parasite
ICHTHYOPHTHIS multifilis, a protozoan which is an obligate parasite
i-antigens, of 48 and 55 kb. Also included in the invention are
i-antigens, of 48 and 55 kb. Also included in the invention are
i-antigens, of binding to the nucleotide sequences for two
antibodies capable of binding to the nucleotide sequences and a method
antipodies capable of binding to the nucleotide sequences.

A composition (containing the i-antigen nucleotide) capable of eliciting
a controlling I multifilis infection in fish. Polynucleotide or protein
controlling I multifilis infection in fish. Polynucleotide or protein
controlling I multifilis infection in fish. Sequences AAA97036-A87042,
preventing I multifilis infection in fish. Sequences and gene
controlling I multifilis infection in fish. Sequences and gene
controlling I multifilis infection in fish. Sequences AAA97036-A87042,
contained to the invention of the invention and peptide sequences.

C AAB25893-B25906 represent 1-antigen protein and peptide sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antigen; i-antigen; ichthyophthiriasis; vaccine;
pase; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               106 CPTGTALDDGVTDVFDRSAAQCVKCKPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTS 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       48kD i-antigen repeat amino acid sequence SEQ ID 10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20.3%; Score 475; DB 21;
100.0%; Pred. No. 3.7e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ilarity 100.0%; Pred. No. 3.7 Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              166 OCVPCQLNKNDSPATAGAQANLATQCSNQ 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  UYGE-) UNIV GEORGIA RES FOUND INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB25863 standard; Protein; 89
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CLARK T G.
DICKERSON H W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0118634.
99US-0122372.
99US-0124905.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              white spot disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 89; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          89 AA;
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02-MAR-1999;
17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-APR-1999;
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This invention relates to novel i-antigen polypeptide sequences.

I antigens or immobilisation antigens ar common to a variety of

I remission to common the common to a variety of

C remission relates and their expression varies in response to

C contrommental stimuli. This invention relates to i-antigens in

C contrommental stimuli. This invention relates to i-antigens in

C controlled two polypeptide and polymucleotide sequences for two

I rantigens, of 48 and 55 kb. Also included in the Invention are

C antibodies capable of binding to the nucleotide sequences and a method

C antibodies capable of binding to the nucleotide sequences and a method

C antibodies capable of binding the i-antigen nucleotide or papelle of eliciting

C antibodies compusing the i-antigen nucleotide or pabelle of

C an immune response in fish is useful for prophylaxis, treatment or for

C antigenic i-antigen polypeptide obtained is also useful for treating

C antigenic i-antigen polypeptide obtained is also useful for treating or

C antigenic i-antigen polypeptide obtained is also useful for treating or

C and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene

C (excluding AAA97060) and AAA97071-A97088 represent primers used in the

C (excluding AAA97060) and AAA97071-A97088 represent primers used in the

C Solation of the I-antigen gene sequences. Sequences. Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish . Disclosure; Figure 5a; 144pp; English. 89 AA; Seguence AAB25861; Query Match 셤 ð ò q

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195 CPTGTVLDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTS 254 Gaps ; 0 20.2%; Score 472; DB 21; Length 89; 100.0%; Pred. No. 6.3e-29; .ive 0; Mismatches 0; Indels 255 QCVPCQINKNDSPATAGAQANLATQCSTQ 283 61 QCVPCQINKNDSPATAGAQANLATQCSTQ 89 AAB25861 standard; Protein; 83 AA. 18-DEC-2000 (first entry) West Local Similarity
watches 89; Conservative

Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control. 48kD i-antigen repeat amino acid sequence SEQ ID 8. (UYGE-) UNIV GEORGIA RES FOUND INC. (CORR ) CORNELL RES FOUND INC. (CLAR/) CLARK T G. 99US-0118634. 99US-0122372. 99US-0124905. 99US-0131121. Ichthyophthirius multifiliis 04-FEB-2000; 2000WO-US02962. WO200046373-A1. 04-FEB-1999; 02-MAR-1999; 17-MAR-1999; 27-APR-1999; 10-AUG-2000 (UYGE-) AAB22861 ID AAB3 XX AAB3 XX AAB3 XX Imm XX Imm XW Wh1 XX ICH PP IO 10 -PP IO 2 -PP IO 2 -PR IO 3 -

This invention relates to novel i-antigen polypeptide sequences.

I antigens or immobilisation antigens ar common to a variety of hymososcomatid cilitates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in controlled their acts of this invention relates to i-antigens in of freshwater fish causing ichthyophthirias or white spot disease. The cof freshwater fish causing ichthyophthirias or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two artibodies capable of binding to the nucleotide sequences and a method for identifying I. multifiliis serotypes using the nucleotide sequences.

A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifiliis infection in fish. Polynucleotide or protein artificial protein of the amplified product encoding an preventing I. multifiliis infection in fish. Sequences AAA9706, and AAA9706, and AAA9706, and AAA9708 represent i antigen gene sequences capable of cacluding AAA97060) and AAA9708 represent primers used in the cacluding AAA97060) and AAA9708 represent primers used in the cacluding AAA97060 and AAA9708 represent primers used in the label substances and peptide sequences.

ABB25893-B25896 represent i antigen gene sequences. Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -Disclosure; Figure 5a; 144pp; English. Ë Lin Dickerson HW, (DICK/) DICKERSON H W. (LINT/) LIN T. WPI; 2000-506071/45. 83 AA; Clark TG, Sequence 

23 CPDGTQTQAGLTDVGAADLGTCVNCRPNFYYNGGAAQGEANGNQPFAANNAARGICVPCQ 82 0; Gaps 19.3%; Score 451; DB 21; Length 83; 100.0%; Pred. No. 2.3e-27; .ive 0; Mismatches 0; Indels Indels ö AAB25865 standard; Protein; 72 AA. 83 INRVGSVTNAGDLATLATQCSTQ 105 61 INRVGSVTNAGDLATLATQCSTQ 83 19.38; 83; Conservative Similarity Query Match Local Matches g οy g

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Immobilisation antigen; 1-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control. 48kD 1-antigen repeat amino acid sequence SEQ ID 12. Ichthyophthirius multifiliis 04-FEB-2000; 2000WO-US02962. 99US-0118634. WO200046373-A1. 04-FEB-1999; 02-MAR-1999; 10-AUG-2000 

18-DEC-2000 (first entry)

AAB25865;

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10-AUG-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 12
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                                                                                                                                                                                                                                                        This invention relates to novel i-antigen polypeptide sequences.

1-antigens or immobilisation antigens ar common to a variety of
1-antigens or immobilisation antigens ar common to a variety of
2. I-antigens or immobilisation antigens ar common to a variety of
2. Invention interpretation and their expression varies in response to
2. Invention includes two polypeptide and polynucleotide sequences for two
2. Invention includes two polypeptide and polynucleotide sequences for two
2. Invention includes two polypeptide and polynucleotide sequences and amethod
3. Invention includes in multifilis sercitypes using the nucleotide sequences.
3. Invention includes in multifilis sercitypes using the nucleotide sequences.
3. Invention in fish is useful for prophylaxis, treatment or for
3. Invention in fish is useful for prophylaxis, treatment or for controlling in multifilis infection in fish. Polynucleotide or protein controlling a portion of the amplified product encoding an antigent i-antigen polypeptide obtained is also useful for treating or antigenic i-antigen for prophylaxis.
3. Preventing I multifilis infection in fish. Sequences AAA97083-A97084
3. Preventing I multifilis infection in fish. Sequences and gene and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene fragments identified in the invention. Sequences AAA97043, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fragments identified in the invention. Sequences AAA97043-A97064 (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolation of the i-antigen gene sequences. Sequences AAB25859-B25889 and AAB25893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                           Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    353 CPAGTVLDDGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVY 412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.2%; Score 379; DB 21; Length 72; 100.0%; Pred. No. 6.5e-22; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   48kD i-antigen repeat amino acid sequence SEQ ID 11.
                                                                                                                                                                                                                                     Disclosure; Figure 5a; 144pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB25864 standard; Protein; 69 AA.
                                             (UYGE-) UNIV GEORGIA RES FOUND INC. (CORR ) CORNELL RES FOUND INC. (CLAR/) CLARR I G.
                                                                                                                              Lin T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chthyophthirius multifiliis
         99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100.
hes 72; Conservative
                                                                                                                               Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                413 AEATOKVOCAST 424
                                                                                      DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200046373-A1.
         17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB25864;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                 TG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                            (CLAR/) (DICK/) I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                 Clark
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This invention relates to novel i-antigen polypeptide sequences.

T-antigens or immobilisation antigens ar common to a variety of rantigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to control of the control of the invention relates to i-antigens in environmental stimuli. This invention relates to i-antigens in controlled realing lichthyophthiriasis or white spot disease. The circhyophthiriasis or disease the control of binding to the nucleotide sequences and a method cirching the i-antigen nucleotide sequences in fish is useful for prophylaxis, treatment or for controlling in multifilis infection in fish. Polynucleotide or protein cantigen polypeptide obtained is also useful for treating or antigence i-antigen polypeptide obtained is also useful for treating or fragments identified in the invention. Sequences AAA37036-A97042, preventing I. multifillis infection in fish. Sequences and gene (excluding AAA37060) and AAA37088 represent i-antigen gene sequences. Sequences AAB25659-B25889 and circle and papersent i-antigen gene sequences. Sequences ABB2589-B35889 and controlling in antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   284 CPTGTAIODGVTLVFSNSSTQCSQCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNTA 343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.1%; Score 376; DB 21; Length 69; 100.0%; Pred. No. 1e-21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DXR; reductoisomerase; enzyme; non-mevalonate isoprenoid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H. influenzae DXR related polypeptide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Pred. w...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABB09437 standard; Protein; 1588 AA.
                                                                                                                                                                               CORNELL RES FOUND INC.
CLARK T G.
                                                                                                                                                                                                                                                                                                                                                                                    Ë
                                                                                                                                                                                                                                                                                                                                                                                    Lin
                                                                                    99US-0122372.
99US-0124905.
99US-0131121.
04-FEB-2000; 2000WO-US02962.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2002 (first entry)
                                                           99US-0118634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100. Matches 69; Conservative
                                                                                                                                                                                                                                                                                                                                                                                       Dickerson HW,
                                                                                                                                                                                                                                                                                            DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          344 TOATQCLTT 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         69 AA;
                                                                                                                                                                                                                                                                                                                                  (LINT/) LIN T.
                                                                                 02-MAR-1999;
17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                           (UYGE-) UNIV
                                                              04-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                          Clark TG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                          (CORR )
(CLAR/)
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The invention relates to modulating an activity of a DXR reductoisomerase enzyme of Haemophilus influenzae, comprising contacting the enzyme with a compound that modulates non-mevalonate isopremoid biosynthesis compound that modulates non-mevalonate isopremoid biosynthesis of menaquinone or ubiquinone. Compounds of the invention act as infected with H. influenzae having DXR reductoisomerase enzyme, e.g. a fineating ear infections, conjunctivitis, meningitis, pneumonia, conjunctivitis, meningitis, pneumonia, conjunctivitis, pneumonia, epiglottitis. The current sequence represents a H. influenzae DXR reductoisomerase enzyme related polypeptide sequence. Once the current sequence contains within it the amino acid sequence of invent in record ABBO9436 (DXR enzyme), but this is broken up by a large consisting the DXR encoding DNA sequence represented as an amino acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24;
                                                                                                                           241..143i - 'region that appears to be accidentally inserted into the sequence, consisting the DXR encoding DNA sequence represented as an amino acid sequence in three letter code"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modulating Haemophilus influenzae DXR reductoisomerase enzyme activity, useful for treating mammals or tissues infected with H. influenzae (e.g. ear infections or pneumonia) by contacting the enzyme with a modulator of its activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        651 TATTGATGCCGTAAAAACTATGGCTCGAAGCTTTTACCAGTAGATAGTGAACATAATGC 710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAACGGGTATTACTGGCAAATAAAGAATCACTGGTAACCTGC------GGACAGCTTTT 650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -----GNOPFAANNAARGIC------VPCQINRVGSVTNAG-DLATLATQC 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            103 STQCPTGTALDDGVTDVFDRSAAQCVKCKPNFYYNGGSPQGEAFGVQVFAAGAAAAGVAA 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              711 TATCTITCAATCATT-----ACGCC-----AGAAGCACAAAAAAATCGGTTT 755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 163 VTSQCVPCQLNKNDSPATAG-----AQANLATQCSNQCPTGTVLDDG-----VTL 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 AVPCPDGTQT-----QAGLTDVGAADL----GT----CVNCRPNFYYNGGAAQGEAN-- 63
             conjunctivitis;
 menaquinone; ubiquinone; virucide; ear infection; conjunctivit:
meningitis; pneumonia; conjunctivitis; bacteraemia; sinusitis;
pleural empyema; endocarditis; epiglottitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels 116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9.4%; Score 219; DB 23; Length 1588; 26.4%; Pred. No. 5.3e-08; Live 14; Mismatches 219; Indels 116
                                                                                                                                                                                                                                                                                                                                                                                                                                       Yan K;
                                                                                                                                                                                                                                                                                                                                                                                                                           Jaworski DD, Payne DJ, Slater-Radosti CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 40-44; 44pp; English.
                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                         (SMIK ) SMITHKLINE BEECHAM CORP. (SMIK ) SMITHKLINE BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                                       09-AUG-2000; 2000US-223909P.
                                                                                                                                                                                                                                                                                                     09-AUG-2001; 2001WO-US24950.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 125; Conservative
                                                                     Haemophilus influenzae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-241698/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1588 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                              WO200211673-A2
                                                                                                                                                                                                                                                                  14-FEB-2002
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                                                                                                          Key
Region
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The present invention describes proteins and their fragments (I) encoded by chromosome 2 of the human malarial parasite, Plasmodium falciparum. Also described are: (1) nucleotide sequences (II) encoding (I); and (2) vaccines against P. falciparum infection comprising (I) or (II).

(I) and (II) are useful for the development of vaccines against P. falciparum infection. (I) and polyclonal antisera or a monoclonal antibody raised to immunogens comprising the sequences of (I), are useful in the detection of infection with P. falciparum. Furthermore, (I) (especially when they are rifins or secreted or membrane proteins) can aid the identification of drugs to treat or prevent P. falciparum infection, or they can be used to identify drug resistance in P. falciparum. Sequencing of the Plasmodium chromosome 2 and the subsequent identification of proteins encoded by it will help to expand our understanding of parasite biology, a process hampered by the complexity of the parasite lightyle, a process hampered by the complexity of the parasite in esistance to drugs and mosquito resistance to insecticides have led to a resurgence of malaria in many
                                                                                                           859 CAÁGCGGTTGCACACCC---TAATTGGTCTATGGGTAAAAATTTCTGT--CGATTCAG 913
208 VENTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQINKNDSP 267
                                                                                                                                                      CPVSKTT----PAHAPG---NTATQATQCLTT----CPAGTVLDDGTSTNFVASATEC-- 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Plasmodium falciparum; chromosome 2; human malaria parasite; vaccine; antimalarial; malaria; protozoacide; infection; insecticide.
                                    816 CCGITACACGCC----ACTTGAA-----CAATTCACCAACATAACACA----GAA
                                                                         268 ATAGAQANLATQCSTQCPTGTAIQDGVTLVFSNSSTQCSQCIANYFFNGNFBAGKSQCLK
                                                                                                                                                                                        CTACAATGATGAATAAGGGCTTGGAATACATTGAGGCTCGCT---GCCTTTTCAATGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Proteins encoded by chromosome 2 of the human malarial parasite, Plasmodium falciparum, useful as antimalarial vaccines and in the diagnosis of P.falciparum infection -
                                                                                                                                                                                                                                                                  971 GIGC-GGAAGAATGGAAGT-TATTATTCATCCACAATCA-ATTATTCATTCTA 1021
                                                                                                                                                                                                                             375 -TKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVYABATQKVQCASTTFA 427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmodium falciparum chromosome 2 related protein SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gardner M, Venter JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 29-33; 577pp; English.
                                                                                                                                                                                                                                                                                                                                                               AAB18144 standard; Protein; 1700 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0107131.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Carucci D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmodium falciparum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HOFF/) HOFFMAN S.
(CARU/) CARUCCI D.
(GARD/) GARDNER M.
(VENT/) VENTER J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-365347/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200025728-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                              07-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                          AAB18144;
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ID AAB1
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Dickerson

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DICKERSON H W.
                                Clark TG,
            (LINT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              δλ
                                                                                                16;
   parts of the world, and there is a pressing need for vaccines and new drugs. AAA70078 to AAA70287 and AAB18144 to AAB18352 represent nucleotide and protein sequences given in the present invention, but which are not specifically mentioned within the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      control
                                                                                                                                                                                                                                                                                        249 VAAVTSQCVPCQINKNDSPATAGAQANLATQCSTQ-----CPTGTAIQDGVTLV-FSNSS 302
                                                                                                                                                                                                                                                                                                                                                    867 IGCGT-----TAG-----AATTTGGAAATATCAATGATTATAGT----A 903
                                                                                                                                                                                                                                                                                                                                                                          363 ISTNFVASATECTKCSAGFFASKTTGFTA----GTDTCTECTKKLTSGATAKVYAEATQ 417
                                                                                                                                                                                                                                                                                                                                                                                            904 AAATTAATAATAAACATAATTTATTGGTAGAAGTGTGTCTTGCAGCCAAATAATGAAGGGG 963
                                                                                                                                                                                                                           703 --AATATGATT-ATAATACTAATGTTACTCATGGTTTTGGTCAAGAGTATCCTTGTGAAA 759
                                                                                                                                                                                                                                                                                                             866
                                                                                                                                        CTGAATTGCATGGTGATTTGTC------AAAAGCAACATATCCAAATGATAAACA 664
                                                                                                                                                             80 PCQINRVGSVTNAGDLATLATQCSTQCPTGTALDDGVTDVFDRSAAQCVKCKPNFYYNGG 139
                                                                                                                    CPDGTQTQAGLTDVGAADLGTCVNCRPNFYINGGAAQG----EANGNQPFAANNAARGICV 79
                                                                                                  Gaps
                                                                                                                                                                                   665 TCCTGAAGGATCAACAGAAATAATCCATG-------CAAACTTC-----
                                                                                                                                                                                                                                                191 CSNQCPT-GTV-LDDGVTLVFNTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAG
                                                                                                                                                                                                                                                                   CGGACATAGTAGAACGTTTTTCTGATACAGAA-----GGAGCACAATGTGATAAGAAAA
                                                                                                                                                                                                                                                                                                            AATAAA-----AGATAATAGTGAAGGAGCTTGCGCTCCATATAGACGATTACATGTA
                                                                                                                                                                                                                                                                                                                                 TQCSQCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTVLDDG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection
                                                                                                  97;
                                                                            DB 21; Length 1700;
                                                                                                  16; Mismatches 202; Indels
                                                                                                                                                                                                        140 SPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNKNDSPATAGAQANLAT-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       55kD i-antigen amino acid repeat sequence SEQ ID 60.
                                                                              9.3%; Score 218.5; DB 2 26.2%; Pred. No. 6.2e-08;
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CORNELL RES FOUND INC.
CLARK I G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             72 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ichthyophthirius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2000; 2000WO-US02962.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB25888 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                             Best Local Similarity 26.2<sup>§</sup>
Matches 112; Conservative
                                                             1700 AA;
                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           964 AATCAAT 970
                                                                                                                                                                                                                                                                                                                                                                                                                        418 KVQCAST 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-FEB-1999;
02-MAR-1999;
17-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB25888;
                                                              Sequence
                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                        303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 14
                                                                                                                            23
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
hymenostomentid ciliates and their expression varies in response to
controlmental stimuli This invention relates to i-antigens in
controlmentar fish causing ichthyophthitiasis or white spot disease. The
cof freshwater fish causing ichthyophthitiasis or white spot disease. The
cof freshwater fish and 55 kD. Also included in the invention are
invention includes two polypeptide and polynucleotide sequences for two
invention includes two polypeptide and polynucleotide sequences of this included in the invention are
contidentifying I. multifillis so included in the invention are
an immune response in fish is useful for prophylaxis; treatment or for
an immune response in fish is useful for prophylaxis; treatment or for
controlling I. multifillis infection in fish. Polynucleotide or protein
controlling I. multifillis infection in fish. Sequences AAA37036-A37042,
and AAA37060, AAA37065 and AAA37089 represent i-antigen genes and gene
fragments identified in the invention. Sequences AAA37036-A37042,
cercluding AAA37060) and AAA37089 represent primers used in the
cercluding AAA37060) and AAA37089 represent primers used in the
cercluding AAA37060 and AAA37089 represent primers used in the
cercluding AAA37060 and AAA37089 represent primers used in the
cercluding AAA37060 and AAA370808 represent primers used in the
cercluding AAA37060 and AAA370808 represent primers used in the
cercluding AAA37060 and AAA370808 represent primers used in the
cercluding AAA37080 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human presentlin enhancer protein pen-1B derived protein SEQ ID NO: 25.
                                                 Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          353 CPAGTVLDDGTSTNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKKLTSGATAKVY 412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use; rat; cow; presenilin enhancer protein; pen; pen-1; pen-1B; pen-2; Aph-2; amyloid beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9.1%; Score 212; DB 21;
13.6%; Pred. No. 4.1e-09;
ve 12; Mismatches 20;
                                                                                                                                                                                                                              Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA014246 standard; Protein; 925 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-MAY-2001; 2001WO-US14648.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-MAY-2000; 2000US-0568942.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            53.6%;
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Best Local Similarity 53.6
Matches 37; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; fruit fly; mouse;
Alzheimer's disease; pen
WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   413 AEATQKVQC 421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | : :||
61 ESAKKNIQC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200185912-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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19;
                                                                                                                                                                                                               The present invention relates to a method of detecting compounds capable of altering the interaction between a presentlin enhancer protein (such as pen.1, pen.18, pen.2 and Aph.2) and presentlin. The inhibition of presentlin activity causes the production of amyloid beta to be reduced and and thus be used in the treatment of Alzheimer's disease. The present exemplification of the invention.
                                                                                                              Presenilin enhancer proteins and polynucleotides useful for modulating presenilin function and screening for an agent that modulates the interaction of the protein to a binding target .
                                                            McGrath GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       205 GAAATATCTGCTGATCTTTGGAGCGTTTGTCTCTGTGTATC--CA-----AGAA 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96 ATLATQC---STQCPTGTALDDGVTDVFDRSAAQCVKCKPNFYXNGGSPQGEA-PGVQVF 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       254 ATGTTCCGATTTGCATATTATAAACTCTTAAAAAAAAGCCAGTGAAGGTTTGAAGAGTATA 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       152 A----AGAAAAGVAAVTSQCVPCQLNKNDSPATAGAQANLATQCSNQCPTGTVLDDGVTL 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       314 AACCCAGGTGAGACAGCACCCTC----TATGGGA--CTGCTGGCCTAT----GTT- 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         208 VENTSATLCVKCRPNFYYNGGSPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQINKNDSP 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         359 -----TCTGGCTTGGGCTTTGGAATCATGAGTGGAGTATTTTCC----- 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           268 ATAGAQANLATQCSTQCPTGTAIQDGVTLVFSNSSTQCS------QCIANYFFNGN 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             398 TITGIGAATACCCTAIC-IGAC-----TCCTIGGGGCCAGGCACAGIGGGCAITCAIGG- 450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               318 FEAGKSQCLKCPVSKTTPAHAPGNTATQATQCLTTCPAGTV-LDDGTSTNFVASATECTK 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAADLGTCVNCR-PNFYYNGGAAQGEANGNQPFAANNAARGICVPCQINRVGSVTNAGDL 95
                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               377 CSAGFFAS-----KTTGFTAGTDICTECTKKLT---SGATAKVYABATQKVQCAS 423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     87;
                                                                                                                                                                                                                                                                                                                                                                                     Query Match

8.9%; Score 208.5; DB 23; Length 925;
Best Local Similarity 25.9%; Pred. No. 1.7e-07;
Matches 108; Conservative 15; Mismatches 207; Indels 87;
                                                            Nicoll SM,
                                                         Ruddy DA,
                                                                                                                                                                              Disclosure; Page 67-70; 78pp; English.
                                                     Ellis MC,
                                                 Curtis DT, Francis GR,
                 (EXEL-) EXELIXIS INC.
                                                                                WPI; 2002-062245/08
                                                                                                                                                                                                                                                                                                                                                        925 AA;
                                                                                                                                                                                                                                                                                                                                                        Seguence
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Search completed: February 11, 2003, 19:46:19 Job time : 35.0571 secs

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